

GenCore version 5.1.4_p5-4578
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OM nucleic - nucleic search, using sw model

Run on: May 3, 2003, 21:10:52 ; Search time 179.041 Seconds
(without alignments)
16512.552 Million cell updates/sec

Title: US-10-027-000-3

Perfect score: 2502
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 746064 seqs, 590810554 residues

Total number of hits satisfying chosen parameters: 1492128

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications NA:*

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- 14: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length DB	ID	Description
1	175.2	7.0	2430	9	US-09-860-846-23
2	175.2	7.0	2430	9	US-09-988-384B-23
3	175.2	7.0	2430	10	US-09-861-289-23
4	175.2	7.0	12441	9	US-09-988-384B-3
5	175.2	7.0	13613	9	US-09-860-846-3
6	175.2	7.0	13613	10	US-09-861-289-3
7	138.6	5.4	2166	9	US-09-808-880-1
8	135.8	5.4	2166	9	US-10-121-032-5
9	135.8	5.4	2166	9	US-10-093-037-5
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11	78.2	3.4	1294	10	US-09-748-033-2
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13	76.2	3.0	2712	10	US-09-748-033-4
14	73.6	2.9	1914	10	US-09-815-242-7960
15	73.2	2.9	2541	10	US-09-476-242-9
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19	71.6	2.9	2541	10	US-09-476-242-12

20	70.8	2.8	1929	9	US-09-899-642-1	Sequence 1, Appl
21	70.8	2.8	13842	9	US-09-860-846-30	Sequence 30, Appl
22	70.8	2.8	13842	9	US-09-988-384B-30	Sequence 30, Appl
23	70.8	2.8	13842	10	US-09-861-289-30	Sequence 30, Appl
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26	70.8	2.8	37988	9	US-09-988-384B-5	Sequence 5, Appl
27	70.8	2.8	2541	10	US-09-476-242-11	Sequence 11, Appl
28	69.8	2.8	88421	9	US-09-976-059-1	Sequence 1, Appl
29	68.8	2.7	1896	9	US-10-124-880-15	Sequence 15, Appl
30	68	2.7	804	9	US-09-773-748-2	Sequence 2, Appl
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35	67	2.7	4509	9	US-10-124-800-5	Sequence 5, Appl
36	67	2.7	4512	9	US-10-124-800-27	Sequence 27, Appl
37	66.6	2.7	905	10	US-09-748-033-7	Sequence 7, Appl
38	66.6	2.7	1965	10	US-09-826-660-26	Sequence 26, Appl
39	66.6	2.7	2322	10	US-09-476-242-20	Sequence 20, Appl
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41	65.4	2.6	390	10	US-09-790-399-7	Sequence 21, Appl
42	65.4	2.6	2310	10	US-09-476-242-21	Sequence 7, Appl
43	65.2	2.6	1248	9	US-09-860-846-7	Sequence 7, Appl
44	65.2	2.6	1248	9	US-09-988-384B-7	Sequence 7, Appl
45	65.2	2.6	1248	10	US-09-861-289-7	Sequence 7, Appl

ALIGNMENTS

RESULT 1
US-09-860-846-23
Sequence 23, Application US/09860846
Patent No. US20020164742A1
GENERAL INFORMATION:
APPLICANT: Sherman, D.H.
APPLICANT: Liu, H.
APPLICANT: Xue, Y.
APPLICANT: Zhao, L.
TITLE OF INVENTION: DNA encoding methymycin and pikromycin
FILE REFERENCE: 600,438051
CURRENT APPLICATION NUMBER: US/09/860,846
CURRENT FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: 09/105,537
PRIOR FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 23
LENGTH: 2430
TYPE: DNA
ORGANISM: Streptomyces venezuelae
US-09-860-846-23

Query Match	7.0%	Score 175.2:	DB 9:	Length 2430:
Best Local Similarity	47.6%	Pred. No. 1.2e+38:		
Matches 1016:	Conservative	0:	Mismatches 968:	Indels 150: Gaps 11:
Db	206	CGCTGGCTTCCACATTAACCAACTGCTCGAAGAGCGAGTAAAGATGATGGCAAG	265	
Qy	206	CGCTGGCTTCCACATTAACCAACTGCTCGAAGAGCGAGTAAAGATGATGGCAAG	265	
Db	353	CCCTGGCCAGCAGCTTGGAGACACATGCGCCAGCAAGTCAAGTATGCGCGCG	412	
Qy	266	AGGCCATGCTTAAGAGTGCAGTGTATCTCGGCGCCGATATCAACATCAACGCTCC	325	
Db	413	ACGTCGCGGCTCAACAGAGACATGCTCGGCGCCGATATCAACATCAACGCTCC	472	
Qy	326	CTCTCGGTGAGAGTGGTTCAGTGTGAGATCCGTCGCGCGCTTGGAG	385	
Db	473	CGCAGCGCGGCGGAACTTCAAGAGCTTCAAGCAGACCCCTGTCTCTCGCGCAGC	532	
Qy	386	CTGCGGCTTCAATCCGAGATTCAGAGCACTGAGTGAAGTCAAGTCAAGCTTTT	445	
Db	533	CGTTCGCGGCAAGTCAAGAGGATCCAGGAGTGTGATGACACAGCGCAACACTTTC	592	

QY	446	TCGCAATGATGAGGAGGACAGGGCAATGATGGTGCAGACATCTGCACGACGGGCTC	505
Db	593	CGGCCACACACCGAGAGAACACCGCTCTCCGTGAACGCCAATGTGCAGCAACAACGC	652
QY	506	TCCGTAATCTACGACACTCCCGTTCCAGATTGCTGTGCAGACATCCACCGGGTGCCT	565
Db	653	TCGCGAGATGAGTTCCCGGGCTTCGAG---GGCTCTCCAAAGCGCGGGCGCTCCT	709
QY	566	TCATGACGGCGGTACATATGGCATATGGCGTGTGTGCACCGAGAACCTAAATATCTTG	625
Db	710	TCATGTGTCCCTACACCGCCTCAACGGGAAGCCGTCCTCGGCACGACGACTCTCA	769
QY	626	ATGGATGCTTTGGAAGGAATGGGGTTGGATGGCTCAATCATGAGGAGCTGTATAGGA	685
Db	770	ACACGTGTGTGGGCACGCAATGGGGCTTCCAGGGCTGGGTATGTCCGATGGCTTGCCA	829
QY	686	CATPACGTAACACAGAACCGCTTGTGGCAGGCTCGACCTCGAGATGCCCGACCTCCAC	745
Db	830	C---CCCGGACCGACGCCCTACCAAGGGCCCTGCACGAGATG-----	873
QY	746	GCTTCCGAGGGAACACTCAAGTTCAACGCTTCGACAGGAAGCCCTTATCCAGCTA	805
Db	874	-----GGGCTGAGGCTTCCCGGACGCTCCGAAAGGCGAGCCCTGCGCCGCGCA	925
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Db	926	ACTTCTTCGGGGA---GGCGCTGAAGACGGCGCTCTGAAACGGCACAGTGTCCGAGCGG	982
QY	866	CGGAAACAGGCCCCGACGACACTGTCAACACACCCCCGAACGGCAGCTCTCTCCGA	925
Db	983	CGGTACCGGTTCCGGCGAGGGATCGTCGGCCAGATGGAAATTCGGCTGTGCTCTCG	1044
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QY	1646	TCGAAATGTCGGTGGCCCTCGCCCAAGGAGCAGACCAAGTCATCTCCGCGGGGCTTA	1705
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QY	1826	GCACCCCGGAGAAATGCTCCCTGGCTGACGCCACGCCCCCTCATCCAGGCTGTGACG	1885
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; Sequence 23, Application US/09988384B			
; Publication No. US20030073824A1			
; GENERAL INFORMATION:			
; APPLICANT: Sherman, D.H.			
; APPLICANT: Liu, H.			
; APPLICANT: Xue, Y.			
; APPLICANT: Zhao, L.			
; TITLE OF INVENTION: DNA encoding methymycin and pikromycin			
; FILE REFERENCE: 600.53GUS1			
; CURRENT APPLICATION NUMBER: US/09/988,384B			
; CURRENT FILING DATE: 2001-11-19			
; PRIOR APPLICATION NUMBER: PCT/US99/14398			
; PRIOR FILING DATE: 1999-06-25			
; PRIOR APPLICATION NUMBER: US 09/105,537			

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RESULT 2
US-09-988-384B-23
; Sequence 23, Application US/09988384B
; Publication No. US20030073824A1
; GENERAL INFORMATION:
; APPLICANT: Sherman, D.H.
; APPLICANT: Liu, H.
; APPLICANT: Xue, Y.
; APPLICANT: Zhao, L.
; TITLE OF INVENTION: DNA encoding methymycin and pikromycin
; FILE REFERENCE: 600.536US1
; CURRENT APPLICATION NUMBER: US/09/988.384B
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: PCT/US99/14398
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,537

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RESULT 3

US-09-861-289-23
 ; Sequence 23, Application US/09861289
 ; Patent No. US2020110897A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Sherman, D. H.
 ; APPLICANT: Liu, H.
 ; APPLICANT: Xue, Y.
 ; APPLICANT: Zhao, L.
 ; TITLE OF INVENTION: DNA encoding methymycin and pikromycin
 ; FILE REFERENCE: 600,438US1
 ; CURRENT APPLICATION NUMBER: US/09/861,289
 ; PRIORITY FILING DATE: 2001-05-18
 ; PRIOR FILING DATE: 1998-06-26
 ; NUMBER OF SEQ ID NOS: 43
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 23
 ; LENGTH: 2430
 ; TYPE: DNA
 ; ORGANISM: Streptomyces venezuelae
 US-09-861-289-23

Query Match 7.0%; Score 175.2; DB 10; Length 2430;

Best Local Similarity 47.6%; Pred. No. 1.2e-38;

Matches 1016; Conservative 0; Mismatches 968; Indels 150; Gaps 11;

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 Db 413 AGGTCGCGCGCTCAACAGGACATGTCTGGGCGCGATGTAACAACATCCGCGTGC 472
 QY 326 CTCTGTGGAGCTGTGGCTTCAGTGTGATGATGATGATGATGATGATGATGATGAT 385
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 Db 1664 TCGGAGAGCGGCTGAG 1715
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Db 5288 TCGCGAAGGCGCTGAGTGTGCGGCGGAGGCGCGTACGCGGCTGCTTTCG----- 5339
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Db 5399 ACAGCTGATCTGCGCTGTGCGGAGACGCCAACCCGACACGATCGTGTGCTCAACACCG 5458
QY 1826 GCAACCCGAGGAGATGCGCTGCGGCTGACGACGCGCGCGCTGATCCAGCGCGCTGATCG 1885
Db 5459 GTTCGTGAGTGTGATGCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5518
QY 1886 GCGGCAACGAGACGCGCAACTCATTTGCCGAGCTGCTTGGGCACTACAAACCCCTCGG 1945
Db 5519 CGGGCCAGGCGGGGCGCGAGGCGCACCGCGCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5578
QY 1946 GCAAGCTGTCTCAGCTTCCG-----AAGCGCTGCAAGAGCAACCCCGCTTTC 1996
Db 5579 GCAAGCTGACGACGACCTTCCGCGCGCGCGGACGACGACGCGCTGCGCGCGGACCGCA 5638
QY 1997 TCAACTTCGCGACCGGCGCGGCGGCGACGCTGACGCGGAGGAGCTGATGCTGCTGCTGCT 2056
Db 5639 CAAGCTACCGCGGCGCTGCGCAACGACGACGACGATACCGGAGGCGCTCCACGCTGGGCTAC 5698
QY 2057 GGTACTACGAGTTTGGCGCAAGAGGAGCTCAATTTCCCTTTGGCGACGCGCTGTCTTCA 2116
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QY 2117 CCACCTTTGCTTTTCAATCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2173
Db 5759 CCTCGTTTCAAGAGAGAGCGCGCGCGCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 5818
QY 2174 CCGTTCGCTGAGAGACCGGCTCCGCGCGCGGCGACAGGTGCGCGCGCTGCTGCTGCTGCT 2233
Db 5819 CGGTTCAGGCTCGCGCAACGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 5878
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Db 5879 GTTCCAGCGCGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 5938
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Db 5939 TCTCGCTCGCGCGCGGCGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 5972

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RESULT 5
US-09-860-846-3

; Sequence 3, Application US/09860846
; Patent No. US20020164742A1
; GENERAL INFORMATION:
; APPLICANT: Sherman, D. H.
; APPLICANT: Liu, H.
; APPLICANT: Xue, Y.
; APPLICANT: Zhao, L.

```

; TITLE OF INVENTION: DNA encoding methymycin and pikromycin
; FILE REFERENCE: 600,438US1
; CURRENT APPLICATION NUMBER: US/09/860,846
; CURRENT FILING DATE: 2001-05-18
; PRIOR APPLICATION NUMBER: 09/105,537
; PRIOR FILING DATE: 1998-06-26
; NUMBER OF SEQ ID NOS: 43
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 13613
; TYPE: DNA
; ORGANISM: Streptomyces venezuelae
; US-09-860-846-3

Query Match      7.0%; Score 175.2; DB 9; Length 13613;
Best Local Similarity 47.6%; Pred. No. 1,9e-38;
Matches 1016; Conservative 0; Mismatches 968; Indels 150; Gaps 11;

QY 206 CGCTCGGTTCCACATTTCACCAACCTGCTGGAAGAGGAGGAGTAAAGTATGAGGCAAG 265
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QY 266 AGCCATCGCTAAGATGCGATGTGATCTCCGCGCGGACTATCAATGACATGCAACGCTCC 325
Db 4724 AGGTCGCGCTCAACCGAGACATGATGCTGGGCGCGATGATGAAACATCCCGGCTGC 4783
QY 326 CTCTCGGTGAGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 385
Db 4784 CGCACGGCGCGGAGACTACGACGACCTTCAACGAGAGACCCCTGCTGCTGCTGCTGCTGCT 4843
QY 386 CTGCGGCTCTCATCCGCGCATTCAGACACTGAGTGCAGAGCTAGCATCAAGCATTTT 445
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QY 446 TGTCAATGATCAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 505
Db 4904 CGGCGCAACCAAGGAGAGCAACCGCTTCTCGGAGGCGCAATGCTGACGAGCAGAGC 4963
QY 506 TCCGTGAATTTAGACACTCCCTTCCAGATTGCTGTGCGAGACTCCACCGGCTGCT 565
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Db 5021 TCATGCTGCTTACACGCGCTCAACGCGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 5080
QY 626 ATGGAGTCTTGAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 685
Db 5081 ACAAGTGTCTGCGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 5140
QY 686 CATACAGTACACAGAGCGCTTGTGCGAGGCGCTGCACTGAGAGTCCCGGAGCTTCAC 745
Db 5141 C---CCGGGAGCGAGCGCATACCAAGGCGCTCGACAGAGAGATG----- 5184
QY 746 GCTTCGAGGAGAAACACTCAAGTTCAAGTCTCAACGAGAAAGCCCTTTATCAGTCA 805
Db 5185 -----GGCGTGTGAGCTCCCGCGAGCTCCGAGAGGCGAGGAGGAGGAGGAGGAGGAG 5236
QY 806 TTGACAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 865
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Db 5414 TCGCGGAGAGCGCGCGGCTCTGCGCAACGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 5473

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QY 1046 GCTCTGCCGCACTAGAGGCGCTACTACGACAGTCACTCCCTTTGAGGGGCTCAGCAAGCAGC 1105
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 QY 1106 TCGAGACGCGCCATCGTACACGCTGCGGCTTACACACCGTCTCTCCATCTTACGCG 1165
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 QY 1166 AGCAGTGCCTACGCGCGCGCGCTCGGCGATGCGCTGAGAGGCTTTCACAGAGCC 1225
 DB 5533 CGCGGGGTGGGGTGGAGCGGTACGTCAGACGGGTAGAGACCTTCGGGACGCG 5652
 QY 1226 CTGGTACCCCTAACCGCGACGACATTTGACGAGCTCTTTCACCAACAGGACATGAC 1285
 DB 5653 ATCCCGGGGGGAACCTCAGC-----CCGGCGTTCAAC 5686
 QY 1286 TGTGTGACTACTACACCCCAAGCGGAGACACGTGTGATCGCGGACATGAGAGCGACGT 1345
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 DB 6086 ACAAGCTATCTCGGCTGTGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 6145
 QY 1826 GCACCGCGGAGAGATGCGCTGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 1885
 DB 6146 GTTGTGCTGCTGATGCG 6205
 QY 1886 GCGGCGAGAGAGCGGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 1945
 DB 6206 CGGGCGAGCG 6265
 QY 1946 GCAAGCTGCTCGAGCTTCCCG-----AAGCGCTGCGAGACGACCGCGCTTC 1996
 DB 6266 GCAAGCTACGAGAGCTTCCCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 6325
 QY 1997 TCAACTTCGCGACGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 2056
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 QY 2057 GGTACTACGAGATTTGCGGACGAGAGCTTCCCGCTTGGCGACGCGCTGCTCTACA 2116
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QY 2117 CCACCTTTGCTTTTCCATCTCTCGTGTCTCACAAGAC---GGCAAGCTGAGCGTGT 2173
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 QY 2234 AGCCCTCCAAAGCG 2293
 DB 6566 GTGCGACCGCGCAAGCTGAGCGCTCCGCGAGCGCAAGAAAGCTCGTGGCTTACGAG 6625
 QY 2294 TCGAAGTCAAGCG 2327
 DB 6626 TCTCGCTCG 6659
 RESULT 6
 US-09-861-289-3
 ; Sequence 3, Application US/09861289
 ; Patent No. US20020110897A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Sherman, D.H.
 ; APPLICANT: Liu, H.
 ; APPLICANT: Xue, Y.
 ; APPLICANT: Zhao, L.
 ; TITLE OF INVENTION: DNA encoding methymycin and plikromycin
 ; FILE REFERENCE: 600 438U1
 ; CURRENT APPLICATION NUMBER: US/09/861,289
 ; PRIOR FILING DATE: 2001-05-18
 ; PRIOR APPLICATION NUMBER: 09/105,537
 ; PRIORITY FILING DATE: 1998-06-26
 ; NUMBER OF SEQ ID NOS: 43
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 3
 ; LENGTH: 13613
 ; TYPE: DNA
 ; ORGANISM: Streptomyces venezuelae
 US-09-861-289-3
 Query Match 7.0%; Score 175.2; DB 10; Length 13613;
 Best Local Similarity 47.6%; Pred. No. 1,9e-38;
 Matches 1016; Conservative 0; Mismatches 968; Indels 150; Gaps 11;
 QY 206 CGCTGCTTCCACATCAACCAACTCTGCTGCAAGAGCGAGTATGATGCGCAAG 265
 DB 4664 CCTGGCCAGACCTTTCAGACGACCATGCGCGACGACTAAGGAGTATGAGCGCG 4723
 QY 266 AGGCAATCGCTAGAGTGGCGCATGTATCTCGCGCGCGCGCGCGCGCGCGCGCG 325
 DB 4724 ACGGTGCG 4783
 QY 326 CTCTGCTGAGCGTGGCTTTCAGTGTGATGAGGATTCCTGCGCGCGCGCGCGCG 385
 DB 4784 CGCAGCG 4843
 QY 386 CTGCGCTTCATCCCGCGGATTCAGACGCTGAGAGTGCAGGTAGATGAGCAACTTT 445
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 QY 446 TGTGCAATGATCAAGAGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 505
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Db 5081 ACAAGCTGCTGGACAGCGAGTGGGGCTTCCAGGGCTGGGTATGTCAGCTGGCTGGCA 5140
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QY 1526 ACAAGTTCAAGATGAGTTGCGGTCCGACCCACCTACACCTCAAGGCGGAGACCATCG 1585
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QY 1646 TCGAAAAGTCCGCTCGCTCGCCAGAGAGACACAGCTCATCTGCGGCGGCGCTTA 1705
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QY 1706 AGCGCAGTGGAGAGACCGGAGGCGCGGAGCGCGGAGCATGAGCTCCCGGCGGTGCTGG 1765

Db 6027 -CTACGAGAGAGGACCGAGGCGCTGACCGCTCGAAGCTGTGCTGCGGAGTACGAGG 6085
QY 1766 ACCAGCTATTTGCGAGAGTGGGCGCGCGGAACCCAAACACCGTGTGATGACAGCG 1825
Db 6086 ACAAGCTGATTCGGCTGTGCGGAGCGCAACCCGAACAGATGATGTCTTCAACACG 6145
QY 1826 GCACCGCGGAGAGATCCCTGCTGACGCGCACCGCGCGCTGATCAGGCTGTACG 1885
Db 6146 GTTCTGCTGATGATCGGCTGTGTCGAAGACCGCGGCTCTGACATGTGTAC 6205
QY 1886 GCGGCAAGAGAGGAGCACTCCATTGCGGAGCTGCTTGTGGGAGTCAACCCCTCG 1945
Db 6206 CGGCGAGGCGGCGGAGGCGCACCGCGCTGTCTGAGGAGTCAACCCCGTTTC 6265
QY 1946 GCAAGCTGCTCCTGAGTTCGCC-----AAGCGCTGACAGACACCCCGCTTTC 1996
Db 6266 GCAAGCTCAGCAAGCTTCCCGGCGCGGCGGAGGAGGAGTCAACCGGCTGCGGAGCG 6325
QY 1997 TCAACTTCGACGAGCGGCGGCGGAGAGCTGACGCGGAGAGAGTCTACGCGGTACA 2056
Db 6326 CAACTACCGCGGCGCTGACCAACGAGAGAGTACCGCGGAGGAGTCACTGCGGTAC 6385
QY 2057 GGTACTAGAGTTTGGCGAGAGAGAGTCAATTTCCTTGGGACAGGCTGCTGCTACA 2116
Db 6386 GCTGCTGACAGAGAGAGTCAAGCGGCTGTCTCCGTTGCGGAGGAGGCTGTGTTACA 6445
QY 2117 CCACTTTTGGCTTTCATCTCTGCTGCTGCTCAAGAGAC---GGCACTGAGCGTGT 2173
Db 6446 CTTGCTTACGCAAGAGGCGCGCGGAGCGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTA 6505
QY 2174 CCTCTCCGTAAGAGACAGCGGCTGCTGCGCGGCGGAGAGTGTGCTGCTGCTGCTA 2233
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QY 2234 AGCCCTTCAGAGGCGGCAAGTTAACCGCCCGCTCAAGAGAGTCAAGGCTTGCAGAG 2293
Db 6566 GTGCGAGCGGAGAGTGTGAGGCTGCTGCGAGGAGGAGGAGTGTGCTGCTGCTGCTG 6625
QY 2294 TCGAAGTCAAGCGCGGAGAGAGAGGCGGTGAC 2327
Db 6626 TCTGCTGCGCGGCGGAGGCGGAGGAGGAGGAGTGTGCTGCTGCTGCTGCTGCTGCTG 6659

RESULT 7
US-09-880-1/c
Sequence 1, Application US/0980880
Publication No. US20030027287A1
GENERAL INFORMATION:
APPLICANT: Belach, Mary C.
APPLICANT: Shah, Sanjay Krishnakant
APPLICANT: McDaniel, Robert
TITLE OF INVENTION: RECOMBINANT OLEANDOLIDE POLYKETIDE SYNTHASE
FILE REFERENCE: 30062-20029.00
CURRENT APPLICATION NUMBER: US/09/808,880
CURRENT FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: US/09/428,517
PRIOR FILING DATE: 1999-10-28
PRIOR APPLICATION NUMBER: 60/120,254
PRIOR FILING DATE: 1999-02-16
PRIOR APPLICATION NUMBER: 60/106,100
PRIOR FILING DATE: 1998-10-29
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patent In Ver. 2.1
SEQ ID NO 1
LENGTH: 50937
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Recombinant DNA
US-09-880-880-1

Query Match

5.5%; Score 138.6; DB 9; Length 50937;

RESULT 10
US-09-790-399-7
: Sequence 7, Application US/09790399
: Patent No. US20020038000A1
: GENERAL INFORMATION:
: APPLICANT: Gold, Larry
: APPLICANT: Therk, Craig
: APPLICANT: Pihlow, David

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: RESULT 11
: US-09-748-033-2
: Sequence 2, Application US/09748033
: Patent No. US20020069431A1
: GENERAL INFORMATION:
: APPLICANT: Broadway, Roxanne M.
: APPLICANT: Congora, Carmenza E.
: TITLE OF INVENTION: EFFECT OF ENOCHITTINASE AND CHIOTOBIOSIDASE AND THEIR
: TITLE OF INVENTION: ENCODING GENES ON PLANT GROWTH AND DEVELOPMENT
: FILE REFERENCE: 19603/3091
: CURRENT APPLICATION NUMBER: US/09/748,033
: CURRENT FILING DATE: 2000-12-22
: PRIOR APPLICATION NUMBER: 60/172,003
: PRIOR FILING DATE: 1999-12-23
: NUMBER OF SEQ ID NOS: 8

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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO: 2
LENGTH: 1294
TYPE: DNA
ORGANISM: Streptomyces albidoflavus
US-09-748-033-2

Query Match 3.1%; Score 78.2; DB 10; Length 1294;
Best Local Similarity 43.8%; Pred. No. 8.2e-12;
Matches 396; Conservative 0; Mismatches 503; Indels 6; Gaps 1;

QY 1224 CCTGTGACCTTACACCGCCAGACATGACAGCTTTCTTACCAAGACGACATGCA 1283
DB 209 CCCCCTTCGACCCCGCGCGCCCTCCACGCCCGCTCCACCGCGCGCGCT 268
QY 1284 CCTGTGACCTTACACCGCCAGACGAGCGGAGACAGCTGTGAGCGGACATGAGGCGAC 1343
DB 269 GACCGCGCGCTTACACCGCGCGCTTCTGCGGCTCCGCGCTCCCTCCACCGCGCGCG 328
QY 1344 GTACACGCGCGAGAGAGCTGACCTTACAGCTGCGCTGCTGCGGCAAGCGCAAA 1403
DB 329 GACCGCGCGAGAGGCG 388
QY 1404 GCGGTAGCTAGACAGCACTGCTGCTGACACGCCACCAAGAGGTCCCGCGGATGC 1463
DB 389 GGTACCGGCTTACGAGCACTTCAACAAAGCGCGCGAGCGCTGCGAGACCTCCCGCGAG 448
QY 1464 CTCTTGGGCTCCGCGCGCGCGCGAGAGCGCGCGCGCTCAATCTGCTCAAGGCGCAACAC 1523
DB 449 GCGCGAGCGCTTACAGCTATCATG-----CGTCTCTTCCGCGCGCGCGCGCGCGCG 502
QY 1524 GTACAGCTTCAAGATGAGTTCGCTGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 1583
DB 503 GCGCGAGATCACTTCAACCTTCAAGCTGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 552
QY 1584 GGTCCCG 1643
DB 563 CCG 622
QY 1644 AATGAAAGTTCGCTGCG 1703
DB 623 CGAGAGGCG 682
QY 1704 TAAACGCGAGTGGAGACCGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 1763
DB 683 CTACGCGCTGATGAGAGTACGCGCTTCAAGCGCGCGCGCGCGCGCGCGCGCGCGCG 742
QY 1764 GAGCGAGCTATTCG 1823
DB 743 CAATCTCAGCTTACATGACCGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 802
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DB 803 GGTCTTCAACATGCG 862
QY 1884 CGCGGCGAGAGAGCG 1943
DB 863 GACGCGCGCTGATCGAAAGCTTCTGACCGCGCGCGCGCGCGCGCGCGCGCGCGCG 922
QY 1944 GGGGAGCTGCTCCCTGAGCTTCCCAAGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 2003
DB 923 CTCGATCTCGGCTGCGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 982
QY 2004 CCGGACGAGGCG 2063
DB 983 GCTGCGCTGATCGAGTGGAGAAAGCTTCTGAGCGCGCGCGCGCGCGCGCGCGCGCG 1042
QY 2064 CGAGTTTCCGAGAGAGCG 2123
DB 1043 CGCTTCCCGAGAGCG 1102
QY 2124 TGCGT 2128

DB 1103 GGAAT 1107

RESULT 12

US-09-748-033-6

Sequence 6, Application US/09748033

Patent No. US20020069431A1

GENERAL INFORMATION:

APPLICANT: Broadway, Roxanne M.

APPLICANT: Gonzora, Carmenza E.

TITLE OF INVENTION: EFFECT OF ENDOCHITINASE AND CHITINOSIDASE AND THEIR

TITLE OF INVENTION: ENCODING GENES ON PLANT GROWTH AND DEVELOPMENT

FILE REFERENCE: 19603/3091

CURRENT FILING DATE: 2000-12-22

PRIOR APPLICATION NUMBER: 60/172,003

PRIOR FILING DATE: 1998-12-23

NUMBER OF SEQ ID NOS: 8

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO: 6

LENGTH: 1107

TYPE: DNA

ORGANISM: Streptomyces albidoflavus

US-09-748-033-6

Query Match 3.0%; Score 76.2; DB 10; Length 1107;
Best Local Similarity 45.7%; Pred. No. 2.8e-11;
Matches 390; Conservative 0; Mismatches 448; Indels 15; Gaps 3;

QY 1129 GTGCGCGCTTACACCGCTTCTCCATTTAGCGAGCACTGCTCCACGCGCGAGCG 1188
DB 118 GCGCGAGATGAGCACTTCCGCTGAGCACTTCCGCGCTTACAGAGCGCGCGCGCGCG 1177
QY 1189 GCTCGCGCGATGCGCTGAGAGGCTTCAACGAGCGCGCGCGCGCGCGCGCGCGCGCG 1248
DB 178 GAGTGGTGGAGCGCGCTCCCGACACTGGAGCAAGCGCGCGCGCGCGCGCGCGCGCG 237
QY 1249 ATTGAGAGCTTCTTTCACCAAGAGAGCACTGCTGAGTATTCACCGCGCGCGCGCG 1308
DB 238 CTCCGAGACTCAAGCG 297
QY 1309 GCGCGAGACAGTGTGATCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 1368
DB 298 ACTGCTGCGCGCGCGCTTCCACCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 357
QY 1369 TACGAGCTGGCGCTGCTGCTGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 428
DB 358 CAGGAGCTGG---TGAGAGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCG 414
QY 1429 GTGCGAGCG 1488
DB 415 TGGGAGTATCCCGAGCG 474
QY 1489 GAGCG 1548
DB 475 AACATGCTCAAGCG 534
QY 1549 TCGGAGCG 1608
DB 535 GCGGAGCG 594
QY 1609 GTGCGCGCGTGAAGGTCTAT-----TGACGAGCGCGCGCGCGCGCGCGCGCG 1662
DB 595 TTGAGATGATCAAGTATGAGTATGAGTATGAGTATGAGTATGAGTATGAGTATGAG 654
QY 1663 CTGCGCGAGAGAGCG 1722
DB 655 ACCG 714
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DB 715 GCG 774

Query Match	2.9%;	Score 73.2;	DB 10;	Length 2541;
Best Local Similarity	43.0%;	Pred. No. 2.4e-10;		
Matches 464;	Conservative 0;	Mismatches 608;	Indels 6;	Gaps 2,

Db 1033 CTGCAAGGCCAGTTGGCAACAGACATCGTTCAAGCAGAGAGGGGGGACCCC 1092
QY 1915 GACGTGCTTTGGGACTACACCCCTGGGCAAGCTGTCCCTCAGTTCCCAAGGCC 1974
Db 1093 GAGATCGTATGACACAGCTTCACTGCGCGCGAGTCTTCTACTGCAACAGACCCAG 1152
QY 1975 CTGCAAGGACACCCCGGCTTCTCAACTTCCGACCCGAGCGGCGCAGCTGTACGGC 2034
Db 1153 CTGTTCAACAGCACCCTGGACACACCCATCGGCCCAACACCAACGACCATCAGC 1212
QY 2035 GAGGACGTCTACGTGCGGTACAGTACTACGAGTTGGCCGACAAAGAGAGTCAATTTCCC 2094
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QY 2095 TTTGGCCACGGCTCTCTTACACCACTTTGGCTTTTCAATCTCTCGGTCTCACAAG 2154
Db 1273 CCATATCCGGGCGGAGATCCGCTGCGAGCAACATCACCGGCTGTGTACCCGCGAC 1332
QY 2155 GACGGCAAGCTGAGGCTGTCCCTCTCCGTGAAGAACACCGGCTCCGCGGCGACAG 2214
Db 1333 GCGGCAAGGAGATCAGCAACACACAGATCTTCCGCCCGCGGCGGAGCATGCGC 1392
QY 2215 GTGGCCAGCTTCTACGTCAAGCCCTCCAGAGGCGCAAGATTAAACCGCCCGTCAAGGAG 2274
Db 1393 GACAACTGGCGCAGAGCTGTACAAGTACAAGGTGTGAAGATCGAGCCCTGGGCGCTG 1452
QY 2275 CTCAGGGCTTGGCAAGGTGCAACTGCAAGCCCGGCGAGAGGCGGTGACAAATCG 2332
Db 1453 GCCCCACCAAGGCCAAGCGCGCGGTGTGAGCGCGAGAAAGCGCGCTGACCTGG 1510

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